

hKCR1
 hKCR1

TMD1
 MAQLGEGYFSAALSCTFLVSCLLPSAFSRALRDEPWDEIFNHLFOAORYCEGRFSLGQNDPMITTLPGLYL
 -----C-----H-----

TMD2 TMD3 TMD4
 VSVGVVKKPASWILGUSEHVCSIGMLRFVNLFSVGNFYLLYLLFEKIQPRKASSIQRIILSTLTAVF
 -----I--F-----*--V-----P-----V-----

TMD5 TMD6
 PTLSTFPMFLYTEAGSVFTLFPAYLMCLYCNHRISALLQFCGPMFROTNITAMAFCAAGHIIAOKCSEAWK
 -----V-----K--P-----V-----NV-----L-----

TMD7
 TELQKKFEERLPPAFGELSELPRVIQPIIHVMSIKNTSMLEPLTTPYVWILLIAPFVFPVVDGGIVVGDR
 -----D-----I-----PA--P-----A-----C-----I--SFL-CA-----I-----

TMD8 TMD9
 SGHEACLIPPOLFYFFCTFAFFCFPHLEPTXVMTPLSLVWIKRRVQFSVITLVSVFLAWKITYV.IKY.LA
 -----L-----S-I-----EGIL-L-V-----A-----

TMD10 TMD11
 DLAHYTFIVWKRVPQRREIVKYLVPAYMFRGAVADSLKSKSIFWNLMFFVCLVASTVPQKLIFFPYFT
 -----A-L-----I-----SI-----P-----I--FIVI-----

TMD12
 LEYIIYRLIMPLPPISELVCELCUYAVVNFITYIFLNKTFQWSDSHDIQRFMW
 -----V-----IT-----T-----S-----I-----I-----ED-Q-----

Figure 1A

10000151.103001

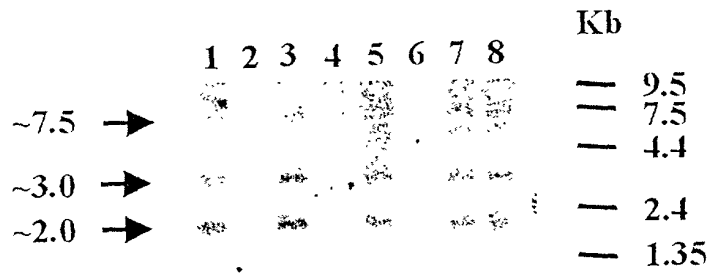


Figure 1B

10000154.103001

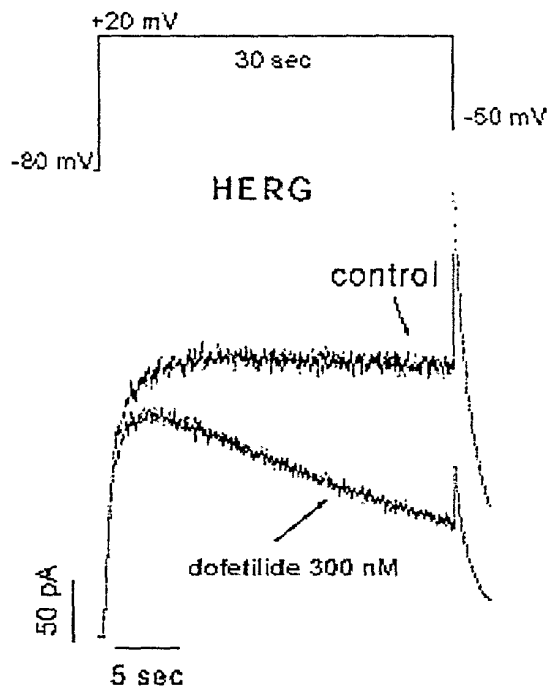


Figure 2A

1000051.103001
T0000T" T5T000T

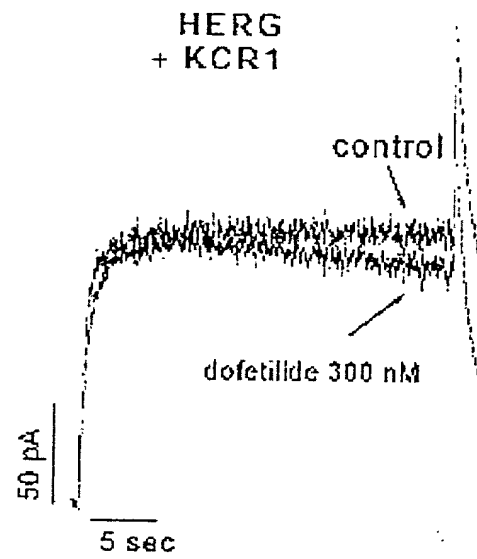


Figure 2B

TOOET 15100001

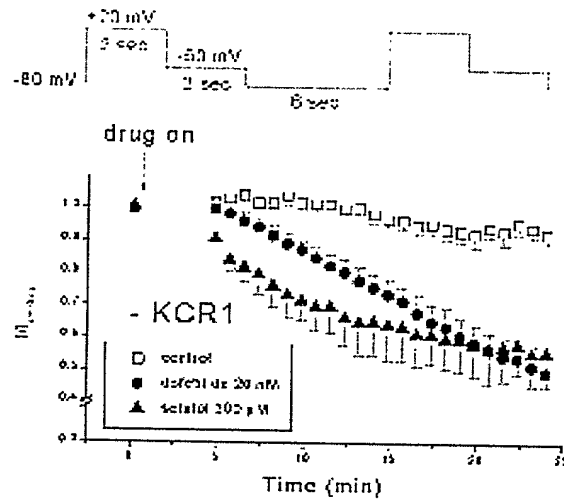


Figure 3A

10000451-103001

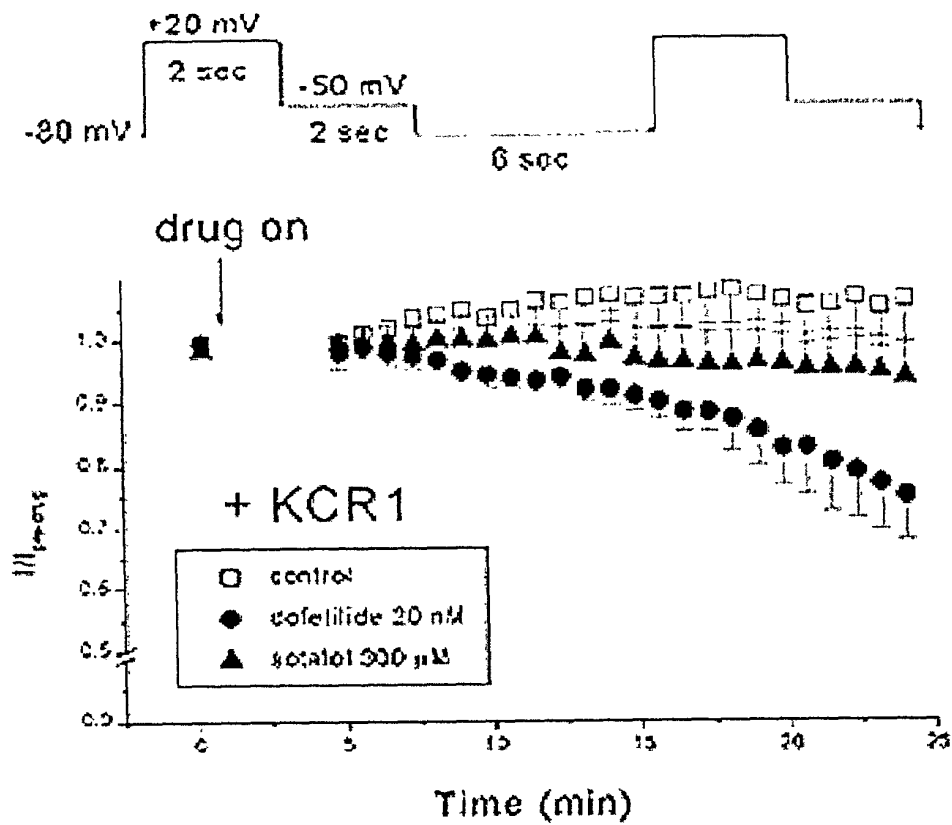


Figure 3B

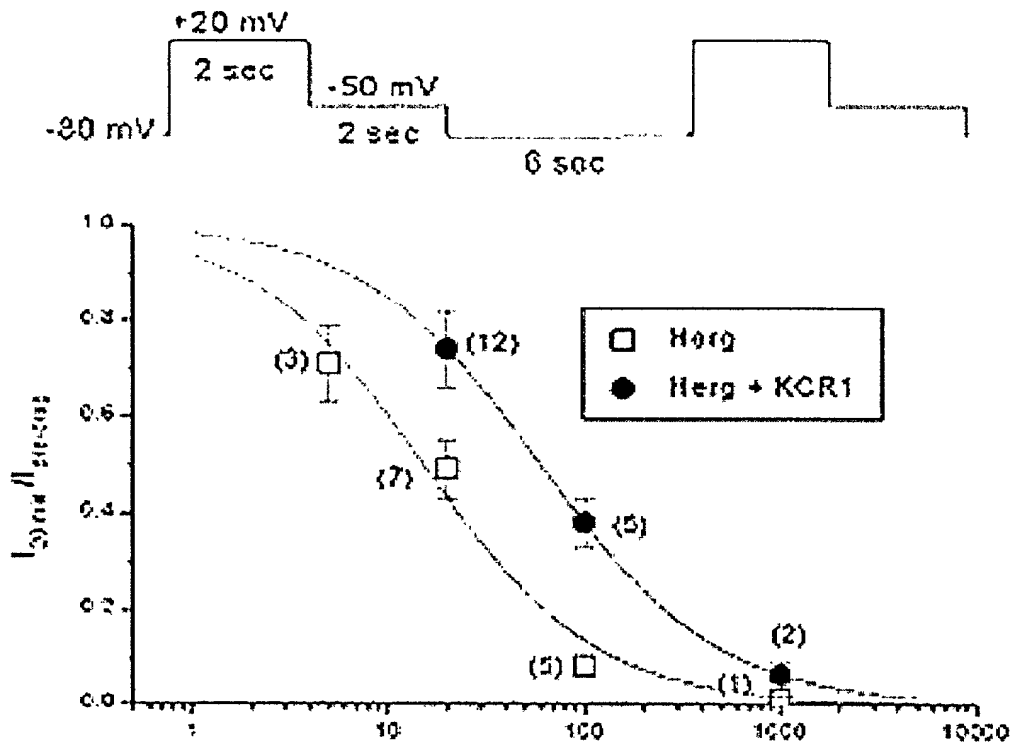


Figure 3C

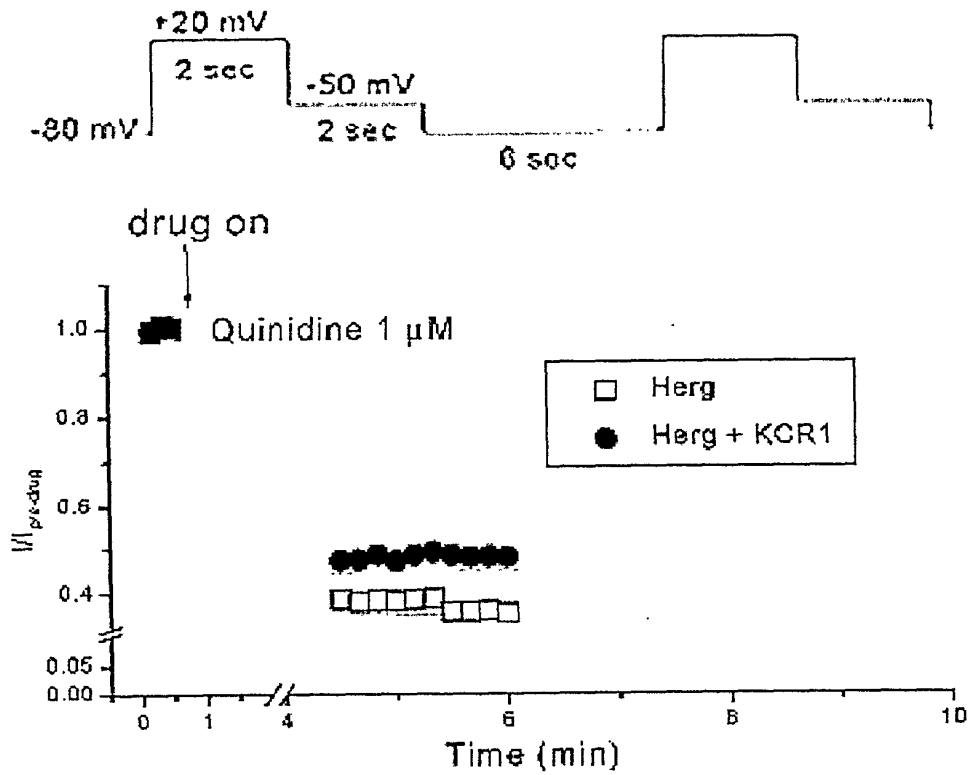


Figure 3D

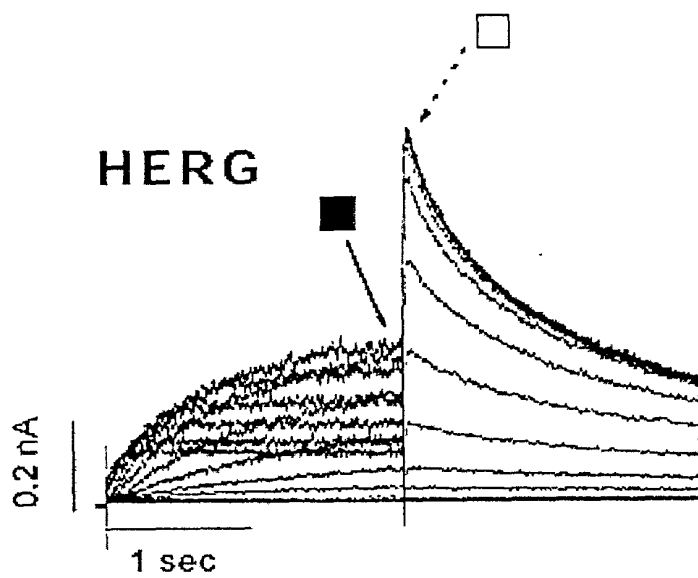


Figure 4A

10000151, 103001

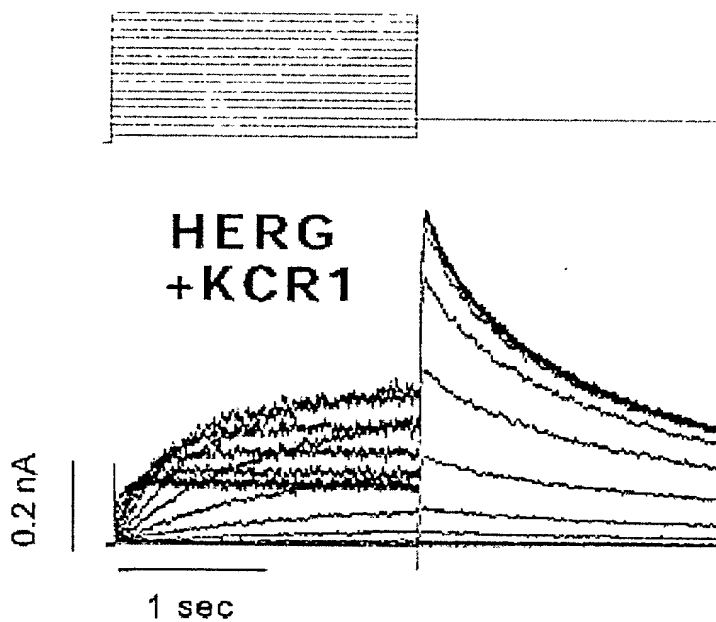


Figure 4B

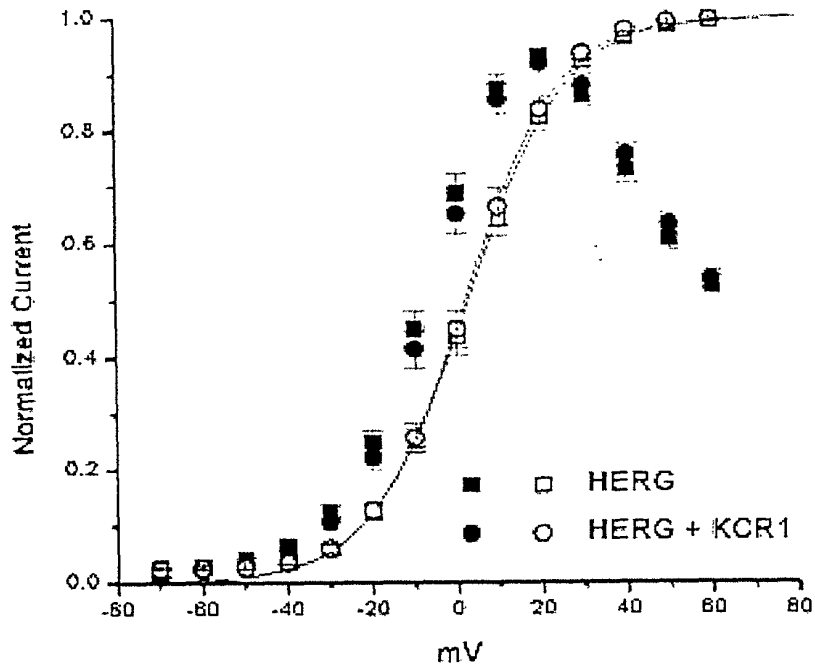


Figure 4C

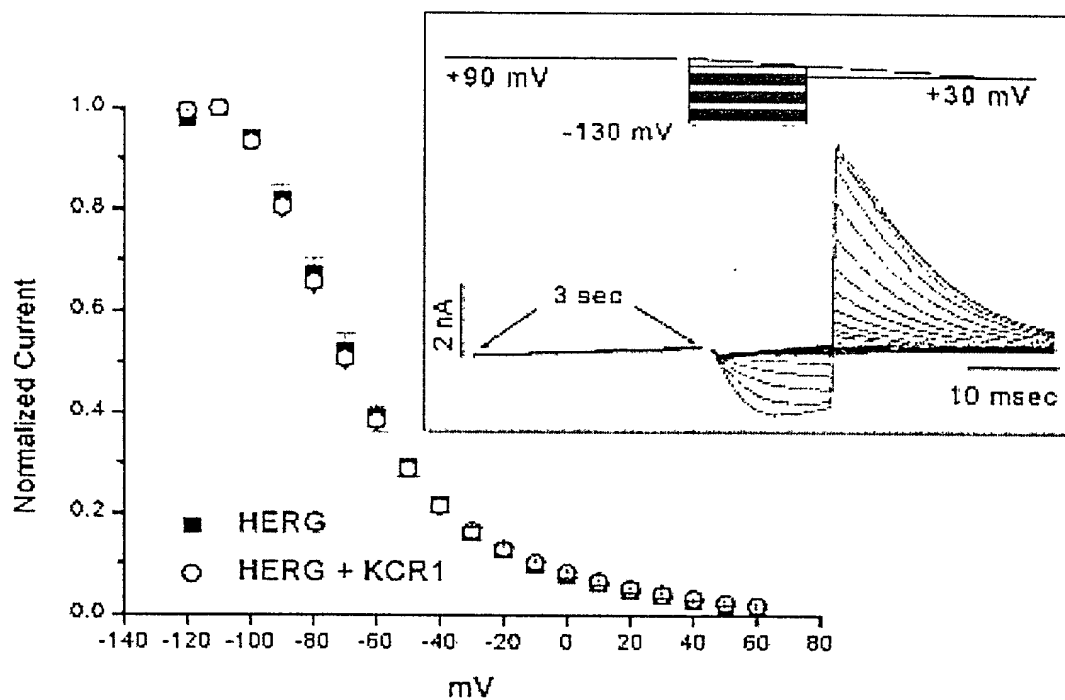


Figure 4D

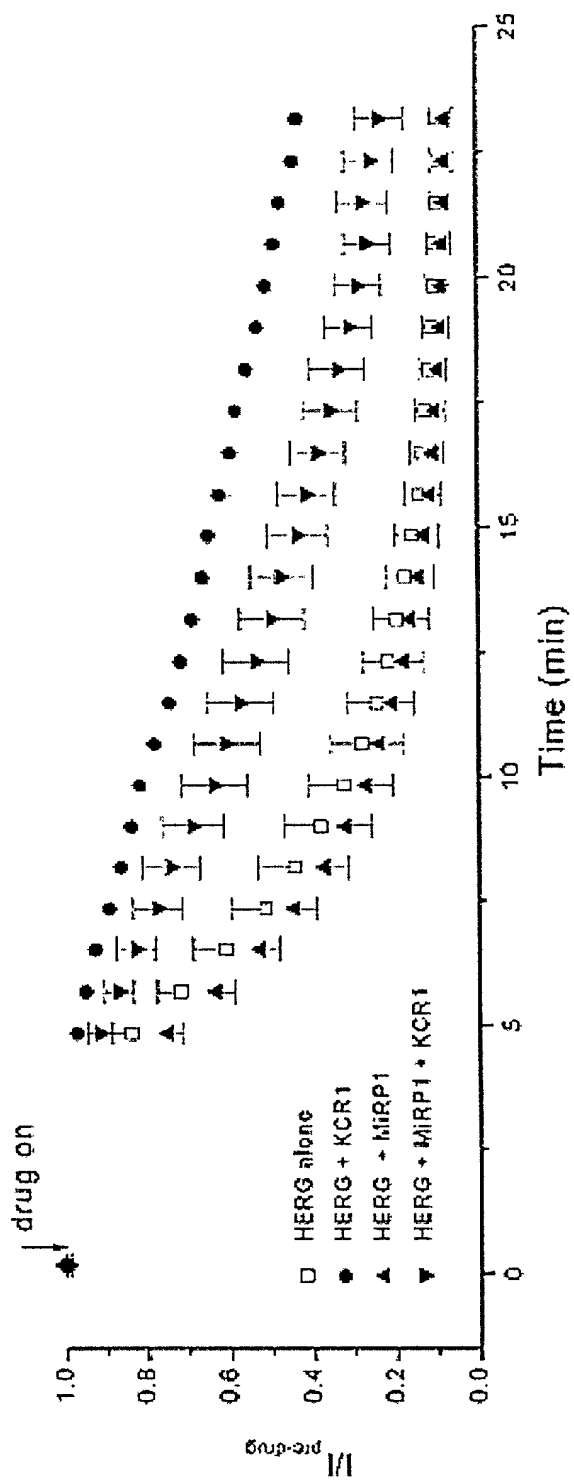


Figure 5A

1000510001

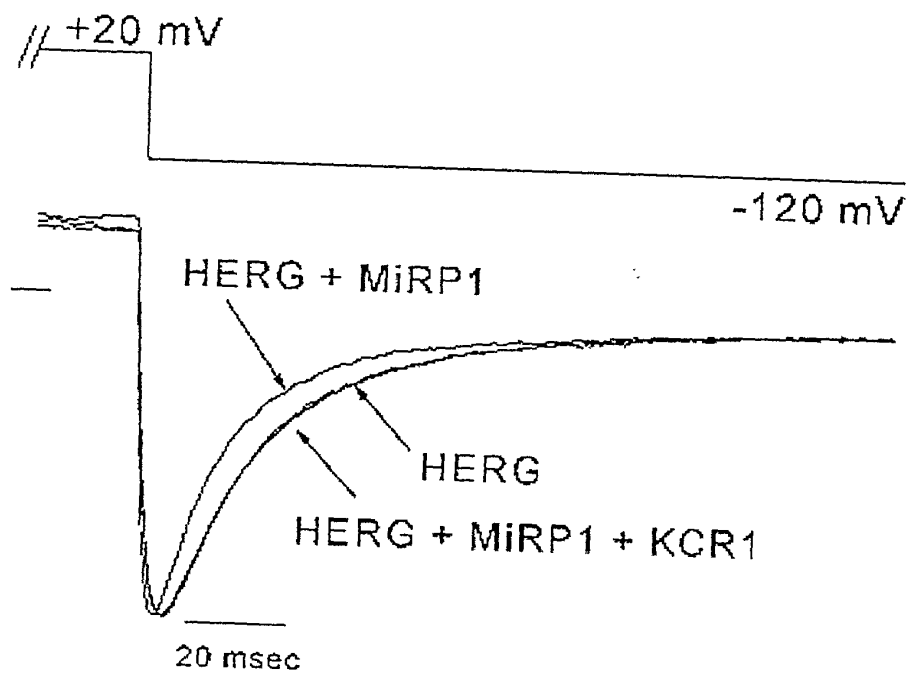


Figure 5B

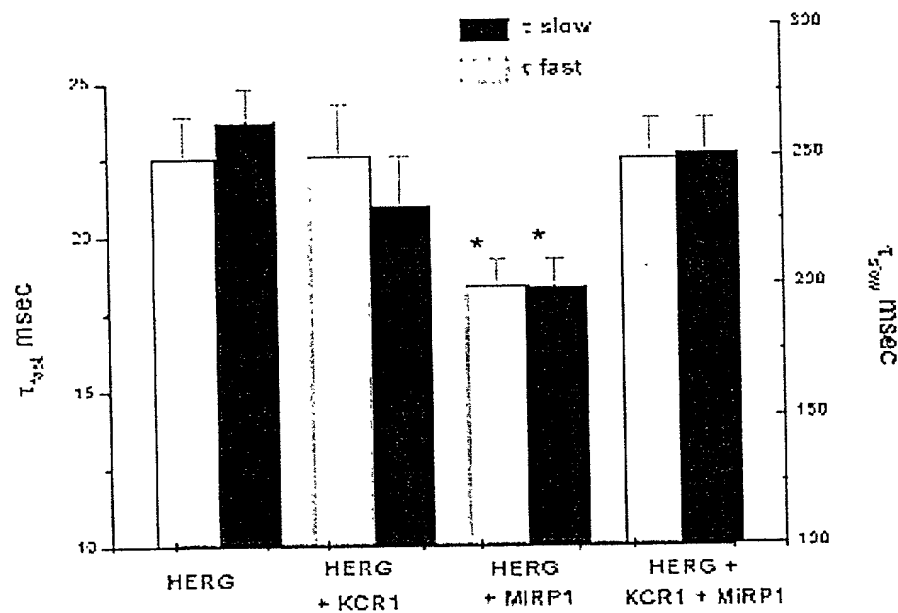


Figure 5C